

AMENDMENTS TO THE CLAIMS

This listing of claims will replace all prior versions and listings of claims in the application:

Listing of claims:

1. (currently amended) A method for SNP typing which comprises the steps of:  
  
simultaneously amplifying a plurality of nucleotide sequences comprising at least one or more sites of single nucleotide polymorphism using genomic DNA whose amount is 10-40 ng per 100 sites and a plurality of primer pairs; and  
  
typing by an INVADER assay or by a TAQMAN PCR method for distinguishing the site(s) of single nucleotide polymorphism of nucleotide sequences amplified in the above amplification step using the amplified nucleotide sequences.
2. (previously presented) The method for SNP typing according to claim 1, wherein said step of amplifying employs a polymerase chain reaction using a hot start method.
3. (original) The method for SNP typing according to claim 1, wherein said step of amplifying employs 50 pairs or more primers.
4. (canceled)

5. (currently amended) The A method for SNP typing ~~according to~~  
~~claim 1~~, which comprises the steps of:  
simultaneously amplifying a plurality of nucleotide sequences  
comprising at least one or more sites of single nucleotide  
polymorphism using genomic DNA whose amount is 10-40 ng and a  
plurality of primer pairs; and  
typing for distinguishing the site(s) of single nucleotide  
polymorphism of nucleotide sequences amplified in the above  
amplification step using the amplified nucleotide sequences  
~~wherein amplification is carried out on from 10 ng to 40 ng of~~  
~~genomic DNA.~~